SEQUENCE LISTING

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<110> Campochiaro, Peter A.
<120> OCULAR GENE THERAPY
<130> OP/4-32696P1
<160> 21
<170> FastSEQ for Windows Version 4.0
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<211> 183
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<213> Human
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Ser Pro Leu Ser Gly Gly Met Arg Gly Ile Arg Gly Ala Asp Phe Gln
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Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala
                             40
                                                 45
Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala
                        55
                                             60
Asp Arg Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe
                    70
                                         75
Pro Ser Trp Glu Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro
                                     90
                                                         95
Gly Ala Arg Ile Phe Ser Phe Asp Gly Lys Asp Val Leu Arg His Pro
                                 105
Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg
                            120
Arg Leu Thr Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser
                        135
Ala Thr Gly Gln Ala Ser Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln
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Ser Ala Ala Ser Cys His His Ala Tyr Ile Val Leu Cys Ile Glu Asn
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Ser Phe Met Thr Ala Ser Lys
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cgggccgtgg
ggctggcggg caccttccgc gccttcctgt cctcgcgcct gcaggacctg
               180
tacagcatcg
tgcgccgtgc cgaccgcgca gccgtgccca tcgtcaacct caaggacgag
ctgctgtttc
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gcacgcatct
               300
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tctcctttga cggcaaggac gtcctgaggc accccacctg gccccagaag 360 atggctcgga ccccaacggg cgcaggctga ccgagagcta ctgtgagacg 420 tggcggacgg aggeteete ggecacggge caggeeteet egetgetggg gggeaggete 480 ctggggcaga gtgccgcgag ctgccatcac gcctacatcg tgctctgcat tgagaacagc ttcatgactg 540 cctccaagta g 551 <210> 3 <211>. 207 <212> PRT <213> Mouse <400> 3 Met Glu Thr Asp Thr Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 10 Gly Ser Thr Gly Asp Ala Ala His Thr His Gln Asp Phe Gln Pro Val 20 25 Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly Met Arg Gly 40 45 Ile Arg Gly Ala Asp Phe Gln Cys Phe Gln Gln Ala Arg Ala Val Gly 55 60 70 75

Leu Ser Gly Thr Phe Arg Ala Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala Asp Arg Gly Ser Val Pro Ile Val Asn 85 90 Leu Lys Asp Glu Val Leu Ser Pro Ser Trp Asp Ser Leu Phe Ser Gly 100 105 110 Ser Gln Gly Gln Leu Gln Pro Gly Ala Arg Ile Phe Ser Phe Asp Gly 115 120 125 Arg Asp Val Leu Arg His Pro Ala Trp Pro Gln Lys Ser Val Trp His 130 135 140 Gly Ser Asp Pro Ser Gly Arg Arg Leu Met Glu Ser Tyr Cys Glu Thr 150 155 Trp Arg Thr Glu Thr Thr Gly Ala Thr Gly Gln Ala Ser Ser Leu Leu

165 170 175

Ser Gly Arg Leu Leu Glu Gln Lys Ala Ala Ser Cys His Asn Ser Tyr
180 185 190

Ile Val Leu Cys Ile Glu Asn Ser Phe Met Thr Ser Phe Ser Lys
195 200 205

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gacgeggece atacteatea ggaettteag ecagtgetee acetggagee
actgaacace 120
cecetgtetg gaggeatgeg tggtateegt ggageagatt tecagtgett
ceageaagee 180
cgageegtgg ggetgteggg cacetteegg gettteetgt ectetagget
geaggatete 240
tatageateg tgegeegtge tgaeeggggg tetgtgeeca tegteaacet
gaaggaegag 300

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gcaacccggg
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gcccgcatct tttcttttga cggcagagat gtcctgagac acccagcctg
                420
gccgcagaag
agcgtatggc acggctcgga ccccagtggg cggaggctga tggagagtta
                480
ctgtgagaca
tggcgaactg aaactactgg ggctacaggt caggcctcct ccctgctgtc
aggcaggctc
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ctggaacaga aagctgcgag ctgccacaac agctacatcg tcctgtgcat
tgagaatagc
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32
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<212> DNA
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Ala Ala
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gcggccgcct atttggagaa agaggtcat
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tttttttc agtgtaaaag gtc
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ctgccagagc cctcccggcc aggcaaagga gaaagaagat ccaggccctc
atggaagctt
               120
ggc
123
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